

[illegible]

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Best Local Similarity   53.3%; Pred. No. 1.3;
Matches    81; Conservative    0; Mismatches    71; Indels    0; Gaps    0;

OY      2003 TTTGTGTGTGTGTGTGTGTGTGTGTGTGTCAGTGTCTCACATGCATGCTGTCACTGG      2062
DB      1132 TGCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT      1191

OY      2063 GTAACAGGCCACCCTCGGACCACTTCATCATACATATTGTAAGTTACTTCCAGCTT      2122
DB      1192 GTCTCTCCCTCTTTTGATTTTGGCCTGGAAATTAATTATTCATATTTCCTGAAAT      1251

Db      2123 CTGGGCATCATTTTGCAATTAATATGCTGTCAAC      2154
      1252 GTGGGTAAACATCTTTAATGAATGAAGTTTTTCTTC      1283


RESULT 5
US-10-664-775-2-COPY/c
GENERAL INFORMATION:
APPLICANT: Simesen, Ruth B
APPLICANT: Pedersen, Anette A
APPLICANT: Fairst, Steffen
APPLICANT: Jensen, Jan J
APPLICANT: Wellguny, Dietmar
TITLE OF INVENTION: Method for Making Recombinant Proteins
FILE REFERENCE: 6448..200-US
CURRENT FILING DATE: 2003-09-17
PRIOR APPLICATION NUMBER: Danish Application PA 2002 01384
PRIOR FILING DATE: 2002-09-20
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 3572
TYPE: DNA
ORGANISM: Baby hamster kidney cell line
US-10-664-775-2-COPY

Query Match          1.2%; Score 29.4; DB 1; Length 3572;
Best Local Similarity 50.3%; Pred. No. 2.5;
Matches    72; Conservative    0; Mismatches    71; Indels    0; Gaps    0;

OY      273 CATCCCCCTGTCTCTCTTCTCCAGAGACTCCAGTCAAGCTGGCGTCATCAAGGGGCCA      332
DB      2639 CGTCCCTCTTCTCTCTCTGCCCCCAATCCCTCCAGCATCAAGGTCCTTTTCCAGTAGAGTCA      2588

OY      333 GACATATCCAACATAGTTCCAGCTTCTGTCCACTGTGACTTTACAGTGTGCTCCCTTTTCA      392
DB      2579 GCTCTTGTGATCAGGTGGCCAAAGTACTGAGTTCAGCTTTAGCATCATTCCTCCCAA      2520

OY      393 GTTACCCAAATCTTGCCCAACAT      415
DB      2519 GAATATCCAGGCTGATCTCTT      2497


RESULT 6
US-10-664-775-2-COPY
GENERAL INFORMATION:
APPLICANT: Simesen, Ruth B
APPLICANT: Pedersen, Anette A
APPLICANT: Fairst, Steffen
APPLICANT: Jensen, Jan J
APPLICANT: Wellguny, Dietmar
TITLE OF INVENTION: Method for Making Recombinant Proteins
FILE REFERENCE: 6448..200-US
CURRENT FILING DATE: 2003-09-17
PRIOR APPLICATION NUMBER: Danish Application PA 2002 01384
PRIOR FILING DATE: 2002-09-20

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/ SEQ ID NO 3
/ LENGTH: 2003
/ TYPE: DNA
/ ORGANISM: Cricetus griseus
US-10-664-775-3-COPY

Query Match          4.3%; Score 25.8; DB 1; Length 2003;
Best Local Similarity 67.9%; Pred. No. 7.2;
Matches 36; Conservative 0; Mismatches 17; Indels 0; Gaps 0;

QY 490 ATTCAATATTTTATTAATGTTAAAAAGCGAGAAAAATTTAAAAAGCGAG 542
DB 1157 ATTCAAGAAAATATGAAATATTAATTCACGCGCCAAAATCAAAAGAGGGG 1105

RESULT 8
US-10-664-775-5-COPY/C
/ Sequence 5, Application US/10664775
/ GENERAL INFORMATION:
/ APPLICANT: Simeesen, Ruth B
/ APPLICANT: Pedersen, Anette A
/ APPLICANT: Faist, Steffen
/ APPLICANT: Jensen, Jan J
/ APPLICANT: Weiguny, Dietmar
/ TITLE OF INVENTION: Method for Making Recombinant Proteins
/ FILE REFERENCE: 6448.200-US
/ CURRENT APPLICATION NUMBER: US/10/664,775
/ PRIOR FILING DATE: 2003-09-17
/ PRIOR APPLICATION NUMBER: Danish Application PA 2002 01384
/ PRIOR FILING DATE: 2002-09-20
/ PRIOR APPLICATION NUMBER: US 60/416,566
/ NUMBER OF SEQ ID NOS: 9
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 5
/ LENGTH: 2267
/ TYPE: DNA
/ ORGANISM: Cricetus griseus
US-10-664-775-5-COPY

Query Match          4.3%; Score 25.8; DB 1; Length 2267;
Best Local Similarity 67.9%; Pred. No. 6.4;
Matches 36; Conservative 0; Mismatches 17; Indels 0; Gaps 0;

QY 490 ATTCAATATTTTATTAATGTTAAAAAGCGAGAAAAATTTAAAAAGCGAG 542
DB 1251 ATTCAAGAAAATATGAAATATTAATTCACGCGCCAAAATCAAAAGAGGGG 1199

RESULT 9
US-10-664-775-4-COPY/C
/ Sequence 4, Application US/10664775
/ GENERAL INFORMATION:
/ APPLICANT: Simeesen, Ruth B
/ APPLICANT: Pedersen, Anette A
/ APPLICANT: Faist, Steffen
/ APPLICANT: Jensen, Jan J
/ APPLICANT: Weiguny, Dietmar
/ TITLE OF INVENTION: Method for Making Recombinant Proteins
/ FILE REFERENCE: 6448.200-US
/ CURRENT APPLICATION NUMBER: US/10/664,775
/ PRIOR FILING DATE: 2003-09-17
/ PRIOR APPLICATION NUMBER: Danish Application PA 2002 01384
/ PRIOR FILING DATE: 2002-09-20
/ PRIOR APPLICATION NUMBER: US 60/416,566
/ NUMBER OF SEQ ID NOS: 9
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 4
/ LENGTH: 2279
/ TYPE: DNA
/ ORGANISM: Cricetus griseus
US-10-664-775-4-COPY
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Query Match          4.3%; Score 25.8; DB 1; Length 2279;
Best Local Similarity 67.9%; Pred. No. 6.4;
Matches 36; Conservative 0; Mismatches 17; Indels 0; Gaps 0;

QY 490 ATTCAATATTTTATTAATGTTAAAAAGCGAGAAAAATTTAAAAAGCGAG 542
DB 1263 ATTCAAGAAAATATGAAATATTAATTCACGCGCCAAAATCAAAAGAGGGG 1211

RESULT 10
US-10-664-775-1-COPY/C
/ Sequence 1, Application US/10664775
/ GENERAL INFORMATION:
/ APPLICANT: Simeesen, Ruth B
/ APPLICANT: Pedersen, Anette A
/ APPLICANT: Faist, Steffen
/ APPLICANT: Jensen, Jan J
/ APPLICANT: Weiguny, Dietmar
/ TITLE OF INVENTION: Method for Making Recombinant Proteins
/ FILE REFERENCE: 6448.200-US
/ CURRENT APPLICATION NUMBER: US/10/664,775
/ PRIOR FILING DATE: 2003-09-17
/ PRIOR APPLICATION NUMBER: Danish Application PA 2002 01384
/ PRIOR FILING DATE: 2002-09-20
/ PRIOR APPLICATION NUMBER: US 60/416,566
/ NUMBER OF SEQ ID NOS: 9
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 1
/ LENGTH: 2715
/ TYPE: DNA
/ ORGANISM: Cricetus griseus
US-10-664-775-1-COPY

Query Match          4.3%; Score 25.8; DB 1; Length 2715;
Best Local Similarity 67.9%; Pred. No. 5.3;
Matches 36; Conservative 0; Mismatches 17; Indels 0; Gaps 0;

QY 490 ATTCAATATTTTATTAATGTTAAAAAGCGAGAAAAATTTAAAAAGCGAG 542
DB 1699 ATTCAAGAAAATATGAAATATTAATTCACGCGCCAAAATCAAAAGAGGGG 1647

Search completed: March 3, 2005, 15:41:38
Job time : 5 secs
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RESULT 2
US-10-664-775-2-COPY
; Sequence 2, Application US/10664775
; GENERAL INFORMATION:
; APPLICANT: Simeesen, Ruth B
; APPLICANT: Pedersen, Anette A
; APPLICANT: Faist, Steffen
; APPLICANT: Jensen, Jan J
; APPLICANT: Wellguy, Dietmar
; TITLE OF INVENTION: Method for Making Recombinant Proteins
; FILE REFERENCE: 6448,200-US
; CURRENT APPLICATION NUMBER: US/10/664,775
; CURRENT FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: Danish Application PA.2002 013847
; PRIOR FILING DATE: 2002-09-20
; PRIOR APPLICATION NUMBER: US 60/416,566
; PRIOR FILING DATE: 2002-10-07
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2
; LENGTH: 3572

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